· C. Kanfman

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### RAW SEQUENCE LISTING PATENT APPLICATION US/08/980,400

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INPUT SET: S32235.raw

This Raw Listing contains the General Information Section and up to the first 5 pages.

1		SEQUENCE LISTING
2 3	(1) Gene	eral Information:
4	(I) Gene	PAITEDED
5	(i) AF	PPLICANT: Sledziewski Ph.D., Andrzej Z ENTERED
6	` '	Bell, Lillian A.
7.		Kindsvogel Ph.D., Wayne R.
8		
9	(ii) Tl	ITLE OF INVENTION: METHODS OF PRODUCING SECRETED RECEPTOR ANALOGS AND BIO
10		·
11	(iii) NU	JMBER OF SEQUENCES: 36
12		
13	• •	DRRESPONDENCE ADDRESS:
1.4	,	(A) ADDRESSEE: Seed and Berry
15		(B) STREET: 6300 Columbia Center, 701 Fifth Avenue
16		(C) CITY: Seattle
17	,	(D) STATE: WA
18		(E) COUNTRY: USA
19	(	(F) ZIP: 98104-7092
20	(\ <b>a</b> c	NADIMED DESPIRE CODY.
21	, ,	OMPUTER READABLE FORM:
22 23		(A) MEDIUM TYPE: Floppy disk (B) COMPUTER: IBM PC compatible
23 24		(B) COMPOTER: 1BM PC COMPACIBLE (C) OPERATING SYSTEM: PC-DOS/MS-DOS
24 25	,	(C) OPERATING SISTEM: PC-DOS/MS-DOS (D) SOFTWARE: PatentIn Release #1.24
26	. '	(D) SOFTWARE: Patentin Release #1.24
27	(vi) CI	JRRENT APPLICATION DATA:
28	, ,	(A) APPLICATION NUMBER: 08/980,400
29	,	(B) FILING DATE:
30	,	(C) CLASSIFICATION:
31	·	( )
32	(vii) PF	RIOR APPLICATION DATA:
33		(A) APPLICATION NUMBER: US 08/477,329
34	Č	(B) FILING DATE: 07-JUN-1995
35		
36	(viii) ATTO	DRNEY/AGENT INFORMATION:
37	(	(A) NAME: Maki, David J.
38		(B) REGISTRATION NUMBER: 31,392
39	(	(C) REFERENCE/DOCKET NUMBER: 990008.446C6
40		
41	, ,	ELECOMMUNICATION INFORMATION:
42		(A) TELEPHONE: 206-622-4900
43	,	(B) TELEFAX: 206-682-6031
44	(	(C) TELEX: 3723836
45		

## RAW SEQUENCE LISTING PATENT APPLICATION US/08/980,400

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INPUT SET: S32235.raw

4/	(2) INFORMATION FOR SEQ ID NO.1:													
48														
49	(i) SEQUENCE CHARACTERISTICS:													
50	(A) LENGTH: 4465 base pairs													
51	(B) TYPE: nucleic acid													
52	(C) STRANDEDNESS: double													
53	(D) TOPOLOGY: linear													
54														
55	(ii) MOLECULE TYPE: cDNA													
56	(II) Moddeodd IIId. edda													
	Add a windowsparant and a													
57	(iii) HYPOTHETICAL: N													
58														
59	(iv) ANTI-SENSE: N													
60														
61	(vi) ORIGINAL SOURCE:													
62	(A) ORGANISM: Homo sapiens													
63	(D) DEVELOPMENTAL STAGE: Adult													
64	(F) TISSUE TYPE: Skin													
65	(G) CELL TYPE: fibroblasts													
66	(6, 6222 11121 12216224616													
67	(vii) IMMEDIATE SOURCE:													
68														
	(B) CLONE: pR-rXl													
69	Alan appendix													
70	(ix) FEATURE:													
71	(A) NAME/KEY: CDS													
72	(B) LOCATION: 3543671													
73	(D) OTHER INFORMATION:													
74														
75														
76	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:													
77														
78	CCCTCAGCCC TGCTGCCCAG CACGAGCCTG TGCTCGCCCT GCCCAACGCA GACAGCCAGA	60												
79														
80	CCCAGGGCGG CCCCTCTGGC GGCTCTGCTC CTCCCGAAGG ATGCTTGGGG AGTGAGGCGA	120												
81														
82	AGCTGGGCGC TCCTCTCCCC TACAGCAGCC CCCTTCCTCC ATCCCTCTGT TCTCCTGAGC	180												
	AGETGGGGGG TECTECCE TACAGEAGEC CECTTECTEC ATECCTEGT TETECTGAGE	100												
83	CEMBOLOGIA COLOGIA CIDA COLOGIA COLOGIA CEMBOLOGIA COLOGIA COL	240												
84	CTTCAGGAGC CTGCACCAGT CCTGCCTGTC CTTCTACTCA GCTGTTACCC ACTCTGGGAC	240												
85														
86	CAGCAGTCTT TCTGATAACT GGGAGAGGGC AGTAAGGAGG ACTTCCTGGA GGGGGTGACT	300												
87														
88	GTCCAGAGCC TGGAACTGTG CCCACACCAG AAGCCATCAG CAGCAAGGAC ACC ATG	356												
89	Met													
90	1													
91	·													
92	CGG CTT CCG GGT GCG ATG CCA GCT CTG GCC CTC AAA GGC GAG CTG CTG	404												
93	Arg Leu Pro Gly Ala Met Pro Ala Leu Ala Leu Lys Gly Glu Leu Leu													
94	5 10 15													
95	•													
96	TTG CTG TCT CTC CTG TTA CTT CTG GAA CCA CAG ATC TCT CAG GGC CTG	452												
97	Leu Leu Ser Leu Leu Leu Leu Glu Pro Gln Ile Ser Gln Gly Leu	772												
98	20 25 30													
99														

# RAW SEQUENCE LISTING PATENT APPLICATION US/08/980,400

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														II	VPUI	SEI:	332233.raw
100	GTC	GTC	ACA	CCC	CCG	GGG	CCA	GAG	CTT	GTC	CTC	AAT	GTC	TCC	AGC	ACC	500
101	Val	Val	Thr	Pro	Pro	Gly	Pro	Glu	Leu	Val	Leu	Asn	Val	Ser	Ser	Thr	
102		35	•				40					45					
103																	
104	TTC	GTT	CTG	ACC	TGC	TCG	GGT	TCA	GCT	CCG	GTG	GTG	TGG	GAA	CGG	ATG	548
105									Ala								
106	50				-	55	-				60		-			65	
107																	
108	TCC	CAG	GAG	ccc	CCA	CAG	GAA	ATG	GCC	AAG	GCC	CAG	GAT	GGC	ACC	TTC	596
109									Ala								-
110	501	<b></b>	014	110	70	01	014	1100	7120	75		<b></b>	p	,	80		
111					, 0					, ,					•		
112	TCC	NGC	СТС	CTC	A C A	CTG	A C C	አአሮ	CTC	λСТ	aca	СПУ	GAC	ACG	GGA	GAA	644
113														_	_	_	011
	Ser	ser	vaı	_	1111	rea	1111	ASII	Leu	IIII.	GLY	Leu	мър		СТУ	GIU	
114				85					90					95			
115	m. a		maa	100	a.a	3 3 m	a.a	maa	aam	<b>aa</b> .	ama	a.a	3.00	C a m	a . a	000	600
116									CGT								692
117	Tyr	Pne		THE	HIS	ASN	ASP		Arg	стХ	Leu	GIU		ASP	GIU	Arg	
118			100					105					110				
119																	7.0
120									GAT								740
121	Lys	_	Leu	Tyr	Ile	Phe		Pro	Asp	Pro	Thr		GTÀ	Phe	Leu	Pro	
122		115					120					125					
123																	
124									TTT								788
125	Asn	Asp	Ala	Glu	Glu	Leu	Phe	Ile	Phe	Leu	Thr	Glu	Ile	Thr	Glu		
126	130					135					140					145	
127																	
128									CCA								836
129	Thr	Ile	Pro	Cys	Arg	Val	Thr	Asp	Pro	Gln	Leu	Val	Val	Thr	Leu	His	
130					150					155					160		
131																	
132	GAG	AAG	AAA	GGG	GAC	GTT	GCA	CTG	CCT	GTC	CCC	TAT	GAT	CAC	CAA	CGT	884
133	Glu	Lys	Lys	Gly	Asp	Val	Ala	Leu	Pro	Val	Pro	Tyr	Asp	His	Gln	Arg	
134				165					170					175			
135																	
136	GGC	TTT	TCT	GGT	ATC	TTT	GAG	GAC	AGA	AGC	TAC	ATC	TGC	AAA	ACC	ACC	932
137	Gly	Phe	Ser	Gly	Ile	Phe	Glu	Asp	Arg	Ser	Tyr	Ile	Cys	Lys	Thr	Thr	
138			180					185					190				
139																	
140	ATT	GGG	GAC	AGG	GAG	GTG	GAT	TCT	GAT	GCC	TAC	TAT	GTC	TAC	AGA	CTC	980
141									Asp								
142		195	•	•			200		_		•	205		-	_		
143																	
144	CAG	GTG	TCA	TCC	ATC	AAC	GTC	TCT	GTG	AAC	GCA	GTG	CAG	ACT	GTG	GTC	1028
145									Val								
146	210					215					220	*				225	
147																	
148	CGC	CAG	GGT	GAG	AAC	АТС	ACC	CTC	ATG	TGC	АТТ	GTG	АТС	GGG	ААТ	GAG	1076
149									Met								20.0
150	9	<b>U 1</b> 11	<b>-</b> -y	<b>01u</b>	230			200		235				~- ]	240		
151					~ ~ ~					255							
152	GTG	GTIC	ልልሮ	ጥጥር	GAG	ጥርር	۵۵۵	ጥልሮ	CCC	רפר	ΔΔΔ	GAA	ΔСΤ	ggg	CGG	СТС	1124
132	010	010	AAC	110	UAU	100	A-A	INC			ллл	UAA	AO I	-555		010	1124

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153 154 155	Val	Val	Asn	Phe 245	Glu	Trp	Thr	Tyr	Pro 250	Arg	Lys	Glu	Ser		D32233.14W
156 157 158 159						GAC Asp									1172
160 161 162 163						CCC Pro									1220
164 165 166 167						GAG Glu 295									1268
168 169 170 171						GTT Val									1316
172 173 174 175						TTT Phe									1364
176 177 178 179						TAC Tyr									1412
180 181 182 183						GAC Asp									1460
184 185 186 187						ACC Thr 375									1508
188 189 190 191						GCT Ala									1556
192 193 194 195						CTC Leu									1604
196 197 198 199						AGT Ser									1652
200 201 202 203						CGG Arg									1700
204 205						AAA Lys									1748

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206	450	•		455			460		IN	<i>PUT</i>	<i>SET:</i> 465	S32235.raw
207 208 209 210			AGT Ser									1796
211 212 213 214 215			GAG Glu 485									1844
216 217 218 219			GAT Asp									1892
220 221 222 223			GAC Asp									1940
224 225 226 227			GTG Val									1988
228 229 230 231			CTT Leu									2036
232 233 234 235			CGA Arg 565									2084
236 237 238 239			TAC Tyr									2132
240 241 242 243			CGG Arg									2180
244 245 246 247			 CAG Gln	 	 	 		 				2228
248 249 250 251			ATG Met									2276
252 253 254 255			AAG Lys 645									2324
256 257 258			CAC His									2372

## **SEQUENCE VERIFICATION REPORT** PATENT APPLICATION *US/08/980,400*

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